

Amendments to the Specification:

Please replace the paragraph beginning at page 11, line 8 with the following amended paragraph:

11,603 clones were isolated. It is known that the baculovirus genome has less than 154 genes/open reading frames. These 11,603 clones should be more than enough to cover all the possible viral genes, the mutations of which would result in nonlytic infection. Among them, 118 mutant clones showed reduction in cell lyses as indicated by the retention of EGFP in cells. In these 118 clones, 20 clones, i.e., A1, A2, A3, C4, 1028, 1044, 1053, 1071, 1081, 1085, 1091, 1094, 3058, 3074, PN8, PN9, PN19, PN23, PN24, and PN121, were further analyzed. Clones 1081 and C4 were deposited, on December 13, 2006, at the China Center for Type Culture Collection, Wuhan, Hubei, 430072, China, where they were given Accession Numbers CCTCC V200610 and CCTCC V200611, respectively.